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ANGIOGENESIS INHIBITORS, COMPOSITIONS CONTAINING THEM AND THEIR USE IN THE TREATMENT OF DISEASES RELATED TO AN ANGIOGENESIS DISORDER

[0001] The present invention concerns the treatment of diseases related to an angiogenesis disorder such as the cancers, arteriosclerosis and diabetes. The invention concerns specific angiogenesis inhibitors able to interfere with the different interactions of TAL-1 likely to be conjugated with vectors such as peptides, particles (liposomes, nanoparticles, etc.) and polymers. The invention also concerns a pharmaceutical composition containing such inhibitors and their use in the treatment of diseases related to angiogenesis, more specifically in the treatment of cancers, arteriosclerosis and diabetes.

[0002] Angiogenesis, process of remodelling pre-existing capillaries, leads to the formation of new vessels [Riseau, Nature 386; 671 (1977)]. In physiological conditions, angiogenesis is closely regulated under the control of a local balance between stimulator (angiogenics) and inhibitor (angiostatics) factors. In the adult, most of the cells of the endothelium (over 99 %) are in the quiescent state. In response to an angiogenic signal, the endothelial cells (ECs) come out of their dormancy and become “activated” to trigger a programme of angiogenesis. These complex processes include the movement of ECs to leave the capillaries and their migration to the angiogenic site, their proliferation and their differentiation to form new capillaries.

[0003] An angiogenesis disorder is observed in certain diseases such as arteriosclerosis, diabetes and tumours [Carmeliet, Nat Med 6; 389 (2000)]. When the tumours reach a critical size creating conditions of hypoxia, the tumour cells give off cytokines, such as VEGF or bFGF, known to initiate or stimulate angiogenesis. Angiogenesis, required for the growth of tumours, is also a main factor in the propagation of malignant cells in the organism (metastases).

[0004] In spite of undeniable progress in the fight against certain cancers, it is necessary to recognise that cancers are progressing globally and that they are a major cause of death. The discovery of new drugs is one of the priorities of medical research. Recent progress in scientific knowledge on tumoral angiogenesis indicates that a treatment based on anti-angiogenesis may provide a new way to control the growth of tumours in cancer patients.

[0005] Several anti-angiogenetic strategies have been developed to inhibit the function of the activator factors (VEGF or bFGF), in particular with neutralising antibodies directed against the cytokine or against its receptor. Other anti-angiogenetic molecules have been derived from the physiological inhibitors present in the haemostatic system, such as plasminogen (angiostatin), collagen (endostatin) or even fibrinogen (alphastatin). These molecules inhibit the adhesion of the ECs to the extracellular matrix required for their morphogenesis.

[0006] Another strategy that has been very little explored to date is to directly target the intracellular events of the endothelial cell in response to angiogenetic stimuli. Therefore, the applicant has demonstrated that TAL-1, a transcription factor from the basic-Helix-Loop-Helix (bHLH) family, specifically modulates the angiogenetic response of the endothelial cells. It controls their migration properties and stimulates their differentiation into capillaries [Lazrak et al., J Cell Sci in press (2004)]. Homologous recombination experiments in the mouse have demonstrated that the tal-1 gene is required for certain stages in haematopoietic and vascular development. Tal-1^{-/-} embryos (not expressing tal-1) present a defective angiogenesis in the yolk sac that seems to reflect an intrinsic defect in the endothelial cells [Visvader et al., Genes Dev 12; 473 (1998)]. In the adult, except of certain haematopoietic progenitors of the bone marrow, TAL-1 protein is expressed in the small vessels in formation, but not in the quiescent mature endothelium [Kallianpur et al., Blood 83; 1200 (1994)]. Significantly, a high level of

expression of TAL-1 is observed in the vasculature of human tumours [Chetty et al., J Pathol 181; 311 (1997)]. TAL-1 is therefore a marker of the angiogenic ECs. Therefore, we targeted TAL-1 to inhibit angiogenesis.

[0007] The applicant developed new molecules able to specifically block the activity of TAL-1 in endothelial cells. The HLH motif of TAL-1 is required for all of the activities of the TAL-1 factor known to date: fixation on DNA or interaction with other nuclear factors, in particular E47 or LMO2 [Hsu et al., Proc. Natl. Acad. Sci 91; 3181 (1994); Vitelli et al., Mol Cell Biol 20; 5330 (2000)] (Figure 1). The applicant developed peptide inhibitors able to enter into competition with the HLH domain of TAL-1. These peptide inhibitors were then coupled with peptide vectors to enable and/or improve their internalisation in the cells.

[0008] The work and results concerning these peptide vectors and their use as vectors to carry active molecules have been described in French patent application no. 97/10297 filed on 08 December 1997.

[0009] In the peptide sequences presented below, the amino acids are represented by their code with one letter, but they may also be represented by their code with three letters according to the following nomenclature.

A	Ala	alanine
C	Cys	cysteine
D	Asp	aspartic acid
E	Glu	glutamic acid
F	Phe	phenylalanine
G	Gly	glycine
H	His	histidine

I	Ile	isoleucine
K	Lys	lysine
L	Leu	leucine
M	Met	methionine
N	Asn	asparagine
P	Pro	proline
Q	Gln	glutamine
R	Arg	arginine
S	Ser	serine
T	Thr	threonine
V	Val	valine
W	Trp	tryptophane
Y	Tyr	tyrosine

[0010] The subject matter of the present invention is a peptide inhibitor (also called peptide molecule) able to interfere with the HLH domain of TAL-1, consisting of or comprising at least 10 successive amino acids and, preferably, at least 15 successive amino acids from the HLH domain of TAL-1 of sequence:

QQNVNGAFAELRKLIPTHTPPDKKLSKNEILRLAMKYINFLA

corresponding to the SEQ ID No. 1 in the listing of sequences in the appendix or an equivalent sequence.

[0011] “Equivalent sequence” refers to the sequence of a known variant of the HLH domain of TAL-1 as well as a sequence presenting one or several substitutions, deletions and/or additions of amino acids with respect to the sequence SEQ ID No. 1, the said substitutions,

deletions and/or additions of amino acids not modifying the activities of the TAL-1 factor thereby obtained as its fixation on DNA or its interaction with other nuclear factors, in particular E47 or LMO 2. The Person Skilled in the Art is aware of different techniques to verify the activity of the TAL-1 transcription factor thereby obtained. In addition, within the framework of the present invention, “equivalent sequence” refers to peptides presenting a post-translational modification and/or a chemical modification, in particular a glycosylation, an amidation, an acylation, an acetylation, a methylation as well as the peptides that carry a protective group. The derivatives of the peptides in the invention may also be those where one or several amino acids are enantiomers, diastereoisomers, natural amino acids of conformation D, rare amino acids in particular hydroxyproline, hydroxylysine, allo-hydroxylysine, 6-N-methyllysine, N-ethylglycine, N-methylglycine, N-ethylasparagine, allo-isoleucine, N-methylisoleucine, N-methylvaline, pyroglutamine, aminobutyric acid and the synthetic amino acids, in particular ornithine, norleucine, norvaline, cyclohexyl-alanine and the omega amino-acids. The derivatives also cover the retropeptides and the retroinverso-peptides, as well as the peptides whose lateral chain of one or several amino acids is substituted by groups that do not modify the anti-microbial activity of the peptides in the invention.

[0012] “Peptide molecule” in the framework of the present invention refers to a molecule consisting of or comprising at least one peptide sequence, the said molecule may also include additional chemical entities other than amino acids.

[0013] Preferably, the inhibitor or peptide molecule, subject matter of the present invention, is chosen from among compound 1 of sequence: QQNVNGAFAELRKLIPTHTPPDKKLSKNEI LRLAMKYINFLA (SEQ ID No. 1 in the list of appended sequences), compound 2 of sequence: VRRIFTNSRERWRQQNVNGAFAELRKLI (SEQ ID No. 2 in the list of appended sequences)

and compound 3 of sequence: PTHPPDKKLSKNEILRLAMKYINFLA (SEQ ID No. 3 in the list of appended sequences) or a sequence equivalent to the said sequences. Therefore, the inhibitor in the present invention consists of or comprises a sequence chosen from among the following sequences:

- QQNVNGAFAELRKLIPTHPPDKKLSKNEILRLAMKYINFLA (SEQ ID No. 1 in the list of appended sequences),
- VRRIFTNSRERWRQQNVNGAFAELRCLI (SEQ ID No. 2 in the list of appended sequences) and
- PTHPPDKKLSKNEILRLAMKYINFLA (SEQ ID No. 3 in the list of appended sequences)

or a sequence equivalent to the said sequences.

[0014] In a preferred embodiment, the inhibitor or peptide molecule, subject matter of the present invention, is associated with a vector. In an advantageous manner, this vector is able to increase the transport of the said inhibitor in the cells and target organs.

[0015] The Person Skilled in the Art is aware of different types of peptide vectors that may be used in the framework of the present invention. Therefore, by way of example and in a non-limiting manner, the vector is chosen from among the group comprising:

- a linear peptide derived from protegrins or tachyplesins [French patent No. 97/10297],
- a linear peptide comprising a domain of transduction such as the domains of transduction of Tat protein of HIV-1 [Fawell et al., Proc. Natl. Acad. Sci 91; 664 (1994); Schwarze et al., Science 285; 1569 (1999) and the domains of transduction derived from

the third helix of Antennapedia [Derossi et al., J. Biol. Chem 269, 10444 (1994); US Patent 5888762],

- particles such as liposomes [Dass and Su. (2001) Drug Deliv. 8:191-213] or nanoparticles [Panyam and Labhasetwar (2003) Adv Drug Deliv Rev. 55:329-47; Douglas et al. (1987) Crit Rev Ther Drug Carrier Syst. 3:233-61],
- polymers such as polyethylene glycol (PEG) [Greenwald et al. (2003) Adv Drug Deliv Rev. 55:217-50].

[0016] The invention in particular envisages as a linear peptide derived from Protegrins, a peptide that complies with the following formula (I):



and as a linear peptide derived from Tachyplesin, a peptide that complies with the following formula (II):



where:

- the identical or different B groups represent an amino acid residue whose lateral chain bears an alkaline group, and
- the identical or different X groups, represent an aliphatic or aromatic amino acid residue

or a fragment of them consisting of a sequence of at least 5 and preferably at least 7 successive amino acids of peptides from formulae (I) or (II).

[0017] The present invention also concerns the use of a vector as defined above to vectorise in the cells, tissues and/or organs, a peptide inhibitor as defined above.

[0018] The bond between the inhibitor or peptide molecule of Tal-1 able to interfere with the different interactions of TAL-1 as defined above and the vector is chosen from among a covalent bond, a hydrophobic bond, an ionic bond, a cleavable bond or a non cleavable bond in physiological media or inside cells.

[0019] This bond may be direct or indirect by means of a linker and carried out by means of a functional group naturally present or introduced either on the vector, or on the inhibitor, or on both. This linker, if present, should be acceptable considering the chemical nature and size of both the vector and inhibitor. By way of example and in a non-limiting manner, we can cite linkers containing alkyl, aryl, aralkyl or peptide groups, esters, aldehydes or alkyl, aryl or aralkyl acids, anhydrid, sulfhydryl or carboxyl groups such as the derivatives of maleymil benzoic acid, maleymil propionic acid and succynimidyl derivatives, the derivatives of cyanogenic bromide or chloride, carbonyldiimidazole, succinimide esters or sulphonic halogenures.

[0020] As functional groups, it is possible to mention: -OH, -SH, -COOH, or -NH₂. Therefore, the inhibitor may be bound by covalent bonds at the level of the N-terminal or C-terminal ends or at the level of the lateral chains of the peptide vector.

[0021] One type of preferred bond between the angiogenesis inhibitor and the vector involves at least one disulphide bridge.

[0022] In a particularly advantageous manner, the inhibitor in the present invention is chosen from among the following two compounds:

- compound 4:



- compound 5:



[0023] The present invention also concerns a pharmaceutical composition comprising, as an active ingredient, at least one peptide inhibitor (or molecule) as defined above, advantageously associated in the said composition with an acceptable vehicle.

[0024] Preferably, the said pharmaceutical composition comes in an appropriate form for parenteral, oral, rectal, nasal, transdermal, pulmonary or central administration. According to the present invention “vehicle” refers to any substance that is added to the inhibitor of the invention to favour its transport, avoid substantial degradation in the said composition and preserve its inhibiting properties. The vehicle is chosen as a function of the type of application listed above the composition.

[0025] The present invention also has for subject matter a method for the treatment of diseases related to an angiogenesis disorder such as the cancers, arteriosclerosis and diabetes, consisting of the administration, to a subject suffering from such a disease, of an efficient quantity of an inhibitor or a composition as described above.

[0026] The invention also concerns the use of an inhibitor or a composition as described previously for the preparation of a drug intended to treat diseases related to angiogenesis, and more specifically diseases related to an angiogenesis disorder, preferably the treatment of cancers, arteriosclerosis and diabetes.

[0027] The invention also concerns the use of a compound able to inhibit the interaction between the HLH domain of TAL-1 and its partner E47 for the preparation of a drug intended to

prevent and/or treat diseases related to angiogenesis and, preferably, the treatment of cancers, arteriosclerosis and diabetes.

[0028] In a first embodiment of the present invention, a compound able to inhibit the interaction between the HLH domain of TAL-1 and its partner E47 is a competitive inhibitor of the HLH domain of TAL-1. Preferably, such a compound is a peptide molecule as defined above.

[0029] In a second embodiment of the present invention, a compound able to inhibit the interaction between the HLH domain of TAL-1 and its partner E47 is a compound able to inhibit the fixation of TAL-1 on its partner E47. In a preferred manner, such a compound is an antibody recognising either an epitope at the level of the HLH domain of TAL-1, or an epitope at the level of the HLH domain of E47.

[0030] The applications and uses considered for the inhibitors (inhibitors or peptide molecules) previously described apply mutates mutandis to the compounds able to inhibit the interaction between the HLH domain of TAL-1 and its partner E47 (methods of treatment, pharmaceutical compositions, etc.).

[0031] Finally, the present invention concerns a method to identify a biologically active compound likely to be used in the prevention and/or treatment of diseases related to angiogenesis and, preferably, the treatment of cancers, arteriosclerosis and diabetes, consisting of detecting the inhibition of the interaction between the HLH domain of TAL-1 and its partner E47 in the presence of the said compound.

[0032] By “biologically active compound”, the present invention more particularly refers to any natural or synthetic chemical compound such as, by way of example but in a non-exhaustive manner, proteins, polypeptides, peptides, aptamers, lipoproteins, polysaccharides, small

molecules, non-peptide molecules, etc. After the identification of the said biologically active compound, it will be easy to test its action in the prevention and/or treatment of diseases related to angiogenesis and, preferably, the treatment of cancers, arteriosclerosis and diabetes using methods familiar to the Person Skilled in the Art.

[0033] The Person Skilled in the Art has different techniques available in order to verify whether the biologically active compound to test is able to inhibit the interaction between the HLH domain of TAL-1 and its partner E47.

[0034] In a first embodiment, the method of the invention comprises the following stages:

- (a) putting into contact the TAL-1 protein (or a fragment of this protein comprising the HLH domain), the transcription factor E47 (or a fragment of this factor comprising the domain that interacts with TAL-1) and the biologically active compound to test,
- (b) immunoprecipitate either the TAL-1 protein (or a fragment of this protein containing the HLH domain), or transcription factor E47 (or a fragment of this factor comprising the domain that interacts with TAL-1),
- (c) if, at stage (b), the TAL-1 protein (or a fragment of this protein containing the HLH domain), is immunoprecipitated, detect in the immunoprecipitate obtained in stage (b), the presence of transcription factor E47 (or a fragment of this factor comprising the domain that interacts with TAL-1),
- (d) if, at stage (b), transcription factor E47 (or a fragment of this factor containing the domain that interacts with TAL-1) is immunoprecipitated, detect in the immunoprecipitate obtained in stage (b), the presence of the TAL-1 protein (or a fragment of this protein comprising the HLH domain),

in case transcription factor E47 (or a fragment of this factor comprising the domain that interacts with TAL-1) is not present in stage (c) or if protein TAL-1 (or a fragment of this protein comprising the HLH domain) is not present in stage (d), the said compound is an agent likely to be used in the prevention and/or treatment of diseases related to angiogenesis and, preferably, the treatment of cancers, arteriosclerosis and diabetes.

[0035] The Person Skilled in the Art has different proteins available that are likely to be used within the framework of the invention method. By way of example, protein E47 that is useable may be the protein from *Gallus gallus* present in Genbank under number CAE30454 or the protein from *Mus musculus* present in Genbank under number AAK18618. In the same way, the TAL-1 protein that may be used is the protein from *Homo sapiens* present in Genbank under number P17542 or the protein from *Mus musculus* present in Genbank under number P22091. In these sequences, the Person Skilled in the Art knows the domains of particular interest within the framework of the present invention.

[0036] In stage (b) of the method of the present invention, the Person Skilled in the Art will know which type of antibody used as a function of the protein to immunoprecipitate [specific antibody of transcription factor E47 (or a fragment of this factor comprising the domain that interacts with TAL-1) or a specific antibody of the TAL-1 protein (or a fragment from this protein comprising the HLH domain)].

[0037] The detection, in stage (c), of the presence, in the immunoprecipitate obtained in stage (b), of transcription factor E47 (or a fragment of this factor comprising the domain that interacts with TAL-1) may be carried out with any technique of the Person Skilled in the Art such as, for example, a Western Blot using a specific antibody for transcription factor E47 (or a fragment of this factor comprising the domain that interacts with TAL-1), an assay of the activity

of transcription factor E47 (or a fragment of this factor comprising the domain that interacts with TAL-1) or a method using a labelled transcription factor E47 (or a fragment of this factor comprising the domain that interacts with TAL-1), the said labelling possibly being radioactive labelling.

[0038] The implementation may be carried out *mutates mutandis* for the TAL-1 protein (or a fragment of this protein comprising the HLH domain) in stage (d).

[0039] In a second embodiment, the method comprises the following stages:

- (a') putting into contact protein TAL-1 (or a fragment of this protein comprising the HLH domain), transcription factor E47 (or a fragment of this factor comprising the domain that interacts with TAL-1) and the biologically active compound to test,
- (b') make migrate on a non-denaturant polyacrylamide gel the mixture obtained in stage (a'),
- (c') visualise the absence or presence of TAL-1 complex (or a fragment of this protein comprising the HLH domain) and E47 (or a fragment of this factor comprising the domain that interacts with TAL-1).

[0040] The absence of this complex in stage (c') reveals the inhibiting effect of the active compound tested and therefore this compound is an agent likely to be used in the prevention and/or treatment of diseases related to angiogenesis and, preferably, the treatment of cancers, arteriosclerosis and diabetes.

[0041] In stage (c') of the method of the invention, different techniques known to the Person Skilled in the Art may be used. By way of example, it is possible to carry out a Western blot using specific antibodies from one of proteins TAL-1 or E47 or their fragments.

[0042] Other advantages and characteristics of the invention will appear upon reading the following examples concerning the preparation of a compound consisting of a TAL-1 inhibitor and a peptide vector as well as on their *in vitro* and *in vivo* effects. Reference will be made to the drawings in the appendix in which:

- figure 1 represents the dimer HLH,
- figure 2 illustrates the *in vitro* inhibition of the interaction of TAL-1 and E47 by the different inhibitors,
- figure 3 illustrates the inhibition of the *in vitro* inhibition of the interaction of TAL-1 and E47 by the inhibitor coupled with a peptide vector,
- figure 4 illustrates the effect of compounds on the survival of HUVEC endothelial cells,
- figure 5 illustrates the *in vitro* effect of one of the compounds in the invention on tubulogenesis of human endothelial cells after 22 hours of culture,
- figure 6 presents the *in vivo* effect of the compounds in the invention on angiogenesis in the mouse by macroscopic analysis of Matrigel implants whether or not complemented with the compounds in the invention,
- figure 7 presents the *in vivo* effect of the compounds in the invention on angiogenesis in the mouse by assay of the haemoglobin contained in the said Matrigel implants.

I – Chemical Synthesis

1) Synthesis of inhibitors of free and vectorised TAL-1

[0043] The peptides were assembled on a solid phase according to a Fmoc strategy, cleaved and deprotected by trifluoroacetic acid, then purified by preparative high pressure

chromatography in inverse phase and lyophilised. Their purity (> 95 %) and their identity were confirmed by analytic HPLC and by mass spectrometry. Peptide sequences: SynB3 (H-RRLSYSRRRF-NH₂; 1394.8 Da, SEQ ID No. 7 in the list of sequences in the appendix), SynB4 (H-AWSFRVSYRGISYRRSR-NH₂; 2145,1 Da, SEQ ID No. 6 in the list of sequences in the appendix), Tal-HLH (H-QQNVNGAFAELRKLIPTHPPDKKLSKNEILRLAMKYINFLA-NH₂, SEQ ID No. 1 in the list of sequences in the appendix).

2) Coupling of inhibitors on peptide vectors


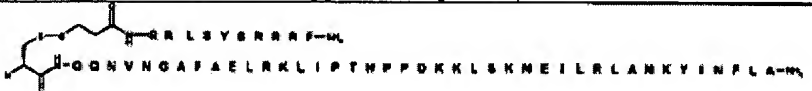
[0044] Activation of the vectors: The peptide vectors SynB3 and SynB4 were treated by SPDP (N-Succinimidyl 3-(2-Pyridylthio)propionate) in Dimethylformamide (DMF) in the presence of DIEA (Diisopropylethylamine). The resulting peptides (2-Pyridylthio)propionyl-SynB3 (1591.8 Da) and (2-Pyridylthio)propionyl-SynB4 (2342.1 Da) were purified by HPLC and lyophilised.

[0045] Preparation of the conjugates: Each activated peptide vector (2-Pyridylthio)propionyl-SynB_x (1eq) was solubilised with C-TalHLH (1eq) in DMF, in the presence of DIEA so as to form a disulphide bridge between the lateral chain of the Cystein of C-TalHLH (Tal-HLH compound synthesised with an additional cysteine) and 3-Mercapto-propionate carried by the vector. The resulting conjugates (Tal-HLH-SynB₃; 6303 Da and Tal-HLH-SynB₄; 7052.6 Da) were precipitated by the addition of ether, then purified by HPLC and lyophilised. The quality control by analytic HPLC at 220 nm and by MALDI-TOF mass spectrometry confirmed the molar masses and determined a purity exceeding 96 %.

II – Compounds tested

[0046] The compounds tested are presented in Table I below.

Table 1

Compound	Name	Sequence
1	Tal-HLH	QQNVNGAFAELRKLIPTHPDKKLSKNEILRLAMKYINFLA (SEQ ID No. 1 in the list of appended sequences)
2	Tal-hel lrb	VRRIFTNSRERWRQQNVNGAFAELRKLI (SEQ ID No. 2 in the list of appended sequences)
3	Tal-hel 2b	PTHPPDKKLSKNEILRLAMKYINFLA (SEQ ID No. 3 in the list of appended sequences)
4	TalHLH-SynB4	 (SEQ ID No. 4 in the list of appended sequences)
5	TalHLH-SynB3	 (SEQ ID No. 5 in the list of appended sequences)
6	SynB4	AWSFRVSYRGISYRRSR (SEQ ID No. 6 in the list of appended sequences)
7	SynB3	RRLSYRRRF (SEQ ID No. 7 in the list of appended sequences)

1) Translation/immunoprecipitation test

[0047] The two proteins TAL-1 and E47 are co-translated *in vitro* using plasmids enabling the transcription and translation of sequences coding in the system TNT-T7-coupled Reticulocyte Lysate, according to the conditions proposed by the supplier (Promega). The translation is carried out in the presence of methionine S³⁵, with or without the addition of peptides dissolved in a 2M solution of urea. An aliquot part (1 µl) of each translation is controlled by electrophoresis and autoradiography of the gel. The addition of up to 0.7 µg of peptide in the reticulocyte lysate (final volume of 25 µl) does not affect the efficacy of the translation of the two proteins.

[0048] After incubation for 30 min. at 37°C, the translation products are immunoprecipitated by a mixture of anti-TAL-1 monoclonal antibodies (BTL 73 + 2TL 136; [Pulford et al., Blood 85; 675 (1995)]) and then analysed by electrophoresis and autoradiography of the gel. The TAL-1-E47 interaction is quantified for each point by calculating the ratio E47/TAL-1 obtained after the autoradiography scan. The intensity of the band corresponding to TAL-1 is used here for

standardisation in all the wells. The interaction in the absence of peptide is arbitrarily established at 100 %.

2) Cytotoxicity Test

[0049] The haematopoietic cells K562 were commercially obtained from ATCC. The cells are seeded with about 10^4 cells per well, 24 h before the addition of the products. They are then at a confluence of 60-80 % on the day of the experiment. The cells are maintained in culture at 37°C in an atmosphere at 95 % humidity and 5 % CO₂ in an OtimMem® medium.

[0050] The cells are incubated with increasing concentrations of the compounds for 48 hours. At the end of the culture time, the MTT (3-(4,5-dimethylthiazole-2-yl)-2,5-diphenyltetrazolium bromide) is added to the wells and the culture plates are then incubated for 4 hours in the oven. The resulting crystalline deposit of formazan is then dissolved with the addition of 200 µl of DMF/SDS. The optical density (OD) is measured at 550 nm (reference 630 nm) using a microplate reader.

[0051] The graphic representation of the percentages of OD of the wells treated as a function of the product concentration is used to determine the LD₅₀. This corresponds to the concentration of the product inhibiting 50 % of the growth.

3) Survival Test of the Endothelial Cells

[0052] The human endothelial cells derived from the vein of the umbilical cord, HUVEC, (Clonetics) are cultivated on dishes coated with gelatine in the complete EMB-2 medium (Clonetics) supplemented with 10 % decomplexed foetal calf serum. The cells are used between passages 3 and 5. The ECV 304 cells (obtained from ATCC) are cultivated in DMEM with 10 % decomplexed foetal calf serum.

[0053] The cells are removed with trypsin and suspended in a medium without serum (OptiMEM, Invitrogen). About 8×10^3 cells in 200 μ l are incubated for 20 min. in the presence of different concentrations of peptides, centrifuged, then seeded in complete EBM2 medium in wells (48-wells plate) coated with collagen I. After 24 hours in the oven, the cell survival is estimated with an MTT test, carried out according to the supplier's (Sigma) recommendations.

III – Results

1) Effect of inhibiting peptides on the *in vitro* interaction of TAL-1/E47

[0054] We chose to test whether the compounds described in Table 1 (Compounds 1, 2 and 3) could affect the formation of hetero dimers TAL-1-E47, mediated by the HLH domain of the two proteins. Compound 1 contains the entire HLH domain of TAL-1 or helix 1- loop-helix 2; compound 2 comprises the basic region and helix 1: compound 3 contains the loop and helix 2 (see figure 1).

[0055] We used a co-immunoprecipitation test for TAL-1 and E47 translated *in vitro* by antibodies directed against one of the proteins.

[0056] Our experiments demonstrate that entire compound Tal-HLH (compound 1) effectively inhibits the interaction of TAL-1 with E47 in a dose-dependant manner (50 to 60 % inhibition with 0.2 μ g; 90 to 100 % with 0.5 μ g) (Figure 2 A and B). The other two inhibiting peptides (compounds 2 and 3) have a lower inhibiting power and do not exceed 50 % with the highest concentrations (Figure 2A). We therefore chose to use inhibiting peptide Tal-HLH (compound 1) for the rest of the experiments.

2) Effect of inhibiting peptides coupled with peptide vectors on the *in vitro* interaction of TAL-1/E47

[0057] We wanted to determine whether the vectorisation or coupling of inhibiting peptide Tal-HLH does not change its inhibiting effect on the *in vitro* interaction of TAL-1 with E47. The

inhibitor was coupled with two peptide vectors (SynB3 and SynB4) via a disulphide bond (compounds 5 and 4, respectively). These two peptide vectors have the ability to increase the transport of molecules through the cell membranes. The results presented in figure 3 confirm that the vectorisation of inhibiting peptide Tal-HLH does not modify its inhibiting effect. It is interesting to note that the inhibitor coupled with vector SynB3 (compound 5) turned out to be more active than the Tal-HLH inhibitor (compound 1).

3) Effect of inhibiting peptides coupled on haematopoietic cells

[0058] The inhibiting effect of vectorised inhibitor was first studied on a haematopoietic line (K562) whose survival requires the activity of TAL-1. The haematopoietic line T (H9) was used as a control since its growth is independent of TAL-1. These two leucemic lines can survive for several days in the absence of growth factors. This can be used to test the effect of peptides in a medium without serum.

[0059] As expected, a significant difference was not observed in the H9 cells (TAL-1 negative), whether the peptide Tal-HLH is vectorised or not (results not shown). However, Tal-HLH peptide had a higher cytotoxic effect on the K562 cells (requiring the action of TAL-1) when vectorised (Table 2). In fact, the LD50, corresponding to the concentration inhibiting 50 % of the growth, is twice as low for the vectorised inhibitor (compound 4) than for the inhibitor alone (compound 1).

Table 2

Compound	Name	LD 50 (μ M)
Compound 1	Tal-HLH	11.5
Compound 2	TalHLH-SynB4	5.7
Compound 3	SynB4	12.3

These results are a first validation that vectorised peptide Tal-HLH (compound 4) is able to specifically inhibit the activity of TAL-1 in the cells.

4) Effect of coupled inhibiting peptides on the survival of endothelial cells

[0060] We also studied the effects of free and vectorised inhibitor on the survival of endothelial cells. Figure 4 shows that compound 1 did not have an effect on the survival of HUVEC cells. However, once coupled (compounds 4 and 5), it specifically affects the survival of the HUVECs cells in a dose-dependant manner. The coupling of the product by a peptide vector was important to produce an effect on survival since the simultaneous addition of inhibitor and non-coupled peptide vector did not have any effect. In the ECV 304 cells used as a negative control, the compounds induced a non specific cell toxicity whether or not coupled (results not shown).

[0061] These experiments demonstrate the specific effect of the TAL-1 inhibitor when it penetrates in the endothelial cells with the peptide vector.

5) Effect of peptides on the *in vitro* tubulogenesis of human endothelial cells (collagen gels in 3-D)

[0062] The HUVEC endothelial cells seeded within a concentrated (1 mg/ml) collagen gel I stop proliferating and, in the presence of activation medium (MDCB131, 1 % foetal calf serum, VEGF 2 ng/ml, bFGF 20 ng/ml and PMA 80 nM), they organise quickly in cell cords to form a primitive network. In 24-48 hours, the aligned cells grow longer, merge together and progressively form pseudo tubules. We tested the effects of vectorised peptide HLH (compound 1) in this *in vitro* system of tubulogenesis of human endothelial cells derived from umbilical cord (HUVECs). Increasing concentrations of vectorised peptide (compound 5) or cargo alone (compound 1) were added both to the collagen solution containing the cells and the activation medium.

[0063] Figure 5 illustrates one of these experiments: vectorised peptide HLH (compound 5) very strongly affects the tubulogenesis between 7 and 10 μ M while the cargo alone (compound

1) does not have an effect on this range of concentrations with respect to the control cultures (5 % DMSO).

6) Effect of peptides on *in vivo* angiogenesis in the mouse (Matrigel plugs)

[0064] The Matrigel plug test consists of a sub-cutaneous injection in the mouse of the Matrigel, an extracellular matrix derived from a murine tumour, able to trigger neovascularisation within the implant. In fact, the endothelial cells of the host, under the influence of the pro-angiogenic factors contained in the Matrigel, migrate towards this “pseudo-tumour” and infiltrate in the Matrigel to organise a network of capillaries between 5 and 7 days.

[0065] In the experiments described above, two sub-cutaneous injections of Matrigel (500 µl) were carried out on both ends of the median line of the abdomen. Just before the injection, the mice were anaesthetised by inhalation of isoflurane (Forène®).

[0066] The Matrigel (BDBiosciences; batch 10403, 13 mg/ml) was complemented by bFGF (500 ng/ml), heparin (60 U/ml) and different peptides SynB3 (compound 7), SynB3-HLH (compound 5), SynB4 (compound 6), SynB4-HLH (compound 4) were added with a final concentration of 20 µM in 5 % DMSO.

[0067] The mice were divided into 3 identical groups of six animals (males C57BL/6, 6 weeks old).

- each mouse in the control group (mice 1 to 6) received two injections of Matrigel containing 5 % DMSO on the left side and a control anti-angiogenic peptide on the right side,
- each mouse in the “SynB3” group (mice 7 to 12) received two injections of Matrigel containing SynB3 on the left side and SynB3-HLH on the right side,

- each mouse in the “SynB4” group (mice 13 to 18) received two injections of Matrigel containing SynB4 on the left side and SynB4-HLH on the right side.

[0068] Six days after the injection, the mice were sacrificed by inhalation of CO₂ and the implants were recovered to assess the angiogenesis (photo and haemoglobin assay). The angiogenesis was quantified by assay of the haemoglobin contained in the Matrigel implants with a solution of Drabkin according to the supplier’s recommendations (Sigma Chemical Co).

[0069] The results presented in Figures 6 (macroscopic analysis) and 7 (haemoglobin assay) demonstrate that the addition of peptide vectors alone did not have a significant effect on angiogenesis. However, HLH peptide conjugated with one of the vectors effectively inhibits neovascularisation in the implants (11 out of 12). This inhibition was higher than that obtained with an inhibitor (Tum) known to inhibit angiogenesis).